1632

Serial Number: 09/909,591

ENTERED

RECEIVED DEC 04 2000 TECH CENTER 1600/2900

CRF Processing Date: 11/6/2000
Edited by: _____
Verified by: _____ (STIC staff)

- [ ] Changed a file from non-ASCII to ASCII
- [ ] Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- [ ] Edited a format error in the Current Application Data section, specifically: _____
- [✓] Edited the Current Application Data section with the actual current number. The number inputted by the applicant was [ ] the prior application data; or [ ] other _____
- [ ] Added the mandatory heading and subheadings for "Current Application Data".
- [ ] Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- [ ] Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- [ ] Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- [ ] Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- [ ] Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- [ ] Inserted colons after headings/subheadings. Headings edited included: _____
- [ ] Deleted extra, invalid, headings used by an applicant, specifically: _____
- [✓] Deleted: [✓] non-ASCII "garbage" at the beginning/end of files; [ ] secretary initials/filename at end of file; [ ] page numbers throughout text; [ ] other invalid text, such as _____
- [ ] Inserted mandatory headings, specifically: _____
- [ ] Corrected an obvious error in the response, specifically: _____
- [ ] Edited identifiers where upper case is used but lower case is required, or vice versa.
- [ ] Corrected an error in the Number of Sequences field, specifically: _____
- [ ] A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- [ ] Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- [ ] Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

```
RAW SEQUENCE LISTING              DATE: 11/13/2000
PATENT APPLICATION:  US/09/509,591   TIME: 10:31:59

Input Set : A:\Pto.amc
Output Set: N:\CRF3\11132000\I509591.raw

SEQUENCE LISTING
 3 (1) GENERAL INFORMATION:
 5       (i) APPLICANT: Little, Andrew
 6                      Lamparski, Henry
 7                      Schuur, Eric
 8                      Henderson, Daniel
10     (ii) TITLE OF INVENTION: ADENOVIRUS VECTORS SPECIFIC FOR CELLS
11                     EXPRESSING ALPHA-FETOPROTEIN AND METHODS OF USE THEREOF
13    (iii) NUMBER OF SEQUENCES: 23
15     (iv) CORRESPONDENCE ADDRESS:
16            (A) ADDRESSEE: MORRISON & FOERSTER
17            (B) STREET: 755 PAGE MILL ROAD
18            (C) CITY: PALO ALTO
19            (D) STATE: CA
20            (E) COUNTRY: USA
21            (F) ZIP: 94304-1018
23      (v) COMPUTER READABLE FORM:
24            (A) MEDIUM TYPE: Floppy disk
25            (B) COMPUTER: IBM PC compatible
26            (C) OPERATING SYSTEM: PC-DOS/MS-DOS
27            (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
29     (vi) CURRENT APPLICATION DATA:
C--> 30            (A) APPLICATION NUMBER: US/09/509,591
C--> 31            (B) FILING DATE: 02-Jun-2000
32            (C) CLASSIFICATION:
34    (vii) PRIOR APPLICATION DATA:
35            (A) APPLICATION NUMBER: PCT/US98/04084
36            (B) FILING DATE: 03-MAR-1998
38   (viii) ATTORNEY/AGENT INFORMATION:
39            (A) NAME: POLIZZI, CATHERINE M.
40            (B) REGISTRATION NUMBER: 40,130
41            (C) REFERENCE/DOCKET NUMBER: 348022000420
43     (ix) TELECOMMUNICATION INFORMATION:
44            (A) TELEPHONE: (650) 813-5600
45            (B) TELEFAX: (650) 494-0792
46            (C) TELEX: 706141 MRSNFOERS SFO
48 (2) INFORMATION FOR SEQ ID NO: 1:
50      (i) SEQUENCE CHARACTERISTICS:
51            (A) LENGTH: 822 base pairs
52            (B) TYPE: nucleic acid
53            (C) STRANDEDNESS: single
54            (D) TOPOLOGY: linear
56     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
58 GCATTGCTGT GAACTCTGTA CTTAGGACTA AACTTTGAGC AATAACACAC ATAGATTGAG      60
60 GATTGTTTGC TGTTAGCATA CAAACTCTGG TTCAAAGCTC CTCTTTATTG CTTGTCTTGG     120
62 AAAATTTGCT GTTCTTCATG GTTTCTCTTT TCACTGCTAT CTATTTTTCT CAACCACTCA     180
64 CATGGCTACA ATAACTGTCT GCAAGCTTAT GATTCCCAAA TATCTATCTC TAGCCTCAAT     240
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RAW SEQUENCE LISTING   DATE: 11/13/2000
PATENT APPLICATION: US/09/509,591   TIME: 10:31:59

Input Set : A:\Pto.amc
Output Set: N:\CRF3\11132000\I509591.raw

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68 TTAAAGACGT TTCAACATAC AAACCGGGGA GTTTTGCCTG GAATGTTTCC TAAAATGTGT    360
70 CCTGTAGCAC ATAGGGTCCT CTTGTTCCTT AAAATCTAAT TACTTTTAGC CCAGTGCTCA    420
72 TCCCACCTAT GGGGAGATGA GAGTGAAAAG GGAGCCTGAT TAATAATTAC ACTAAGTCAA    480
74 TAGGCATAGA GCCAGGACTG TTTGGGTAAA CTGGTCACTT TATCTTAAAC TAAATATATC    540
76 CAAAACTGAA CATGTACTTA GTTACTAAGT CTTTGACTTT ATCTCATTCA TACCACTCAG    600
78 CTTTATCCAG GCCACTTATG AGCTCTGTGT CCTTGAACAT AAAATACAAA TAACCGCTAT    660
80 GCTGTTAATT ATTGGCAAAT GTCCCATTTT CAACCTAAGG AAATACCATA AGTAACAGA     720
82 TATACCAACA AAAGGTTACT AGTTAACAGG CATTGCCTGA AAAGAGTATA AAAGAATTTC    780
84 AGCATGATTT TCCATATTGT GCTTCCACCA CTGCCAATAA CA                       822
86 (2) INFORMATION FOR SEQ ID NO: 2:
88      (i) SEQUENCE CHARACTERISTICS:
89          (A) LENGTH: 5224 base pairs
90          (B) TYPE: nucleic acid
91          (C) STRANDEDNESS: single
92          (D) TOPOLOGY: linear
94      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
96 GAATTCTTAG AAATATGGGG GTAGGGGTGG TGGTGGTAAT TCTGTTTTCA CCCCATAGGT     60
98 GAGATAAGCA TTGGGTTAAA TGTGCTTTCA CACACACATC ACATTTCATA AGAATTAAGG    120
100 AACAGACTAT GGGCTGGAGG ACTTTGAGGA TGTCTGTCTC ATAACACTTG GGTTGTATCT    180
102 GTTCTATGGG GCTTGTTTTA AGCTTGGCAA CTTGCAACAG GGTTCACTGA CTTTCTCCCC    240
104 AAGCCCAAGG TACTGTCCTC TTTTCATATC TGTTTTGGGG CCTCTGGGGC TTGAATATCT    300
106 GAGAAAATAT AAACATTTCA ATAATGTTCT GTGGTGAGAT GAGTATGAGA GATGTGTCAT    360
108 TCATTTGTAT CAATGAATGA ATGAGGACAA TTAGTGTATA AATCCTTAGT ACAACAATCT    420
110 GAGGGTAGGG GTGGTACTAT TCAATTTCTA TTTATAAAGA TACTTATTTC TATTTATTTA    480
112 TGCTTGTGAC AAATGTTTTG TTCGGGACCA CAGGAATCAC AAAGATGAGT CTTTGAATTT    540
114 AAGAAGTTAA TGGTCCAGGA ATAATTACAT AGCTTACAAA TGACTATGAT ATACCATCAA    600
116 ACAAGAGGTT CCATGAGAAA ATAATCTGAA AGGTTTAATA AGTTGTCAAA GGTGAGAGGG    660
118 CTCTTCTCTA GCTAGAGACT AATCAGAAAT ACATTCAGGG ATAATTATTT GAATAGACCT    720
120 TAAGGGTTGG GTACATTTTG TTCAAGCATT GATGGAGAAG GAGAGTGAAT ATTTGAAAAC    780
122 ATTTTCAACT AACCAACCAC CCAATCCAAC AAACAAAAAA TGAAAAGAAT CTCAGAAACA    840
124 GTGAGATAAG AGAAGGAATT TTCTCACAAC CCACACGTAT AGCTCAACTG CTCTGAAGAA    900
126 GTATATATCT AATATTTAAC ACTAACATCA TGCTAATAAT GATAATAATT ACTGTCATTT    960
128 TTTAATGTCT ATAAGTACCA GGCATTTAGA AGATATTATT CCATTTATAT ATCAAAATAA   1020
130 ACTTGAGGGG ATAGATCATT TTCATGATAT ATGAGAAATA TTAAAAACAG ATTGAATTAT   1080
132 TTGCCTGTCA TACAGCTAAT AATTGACCAT AAGACAATTA GATTTAAATT AGTTTTGAAT   1140
134 CTTTCTAATA CCAAAGTTCA GTTTACTGTT CCATGTTGCT TCTGAGTGGC TTCACAGACT   1200
136 TATGAAAAAG TAAACGGAAT CAGAATTACA TCAATGCAAA AGCATTGCTG TGAACTCTGT   1260
138 ACTTAGGACT AAACTTTGAG CAATAACACA CATAGATTGA GGATTGTTTG CTGTTAGCAT   1320
140 ACAAACTCTG GTTCAAAGCT CCTCTTTATT GCTTGTCTTG GAAAATTTGC TGTTCTTCAT   1380
142 GGTTTCTCTT TTCACTGCTA TCTATTTTTC TCAACCACTC ACATGGCTAC AATAACTGTC   1440
144 TGCAAGCTTA TGATTCCCAA ATATCTATCT CTAGCCTCAA TCTTGTTCCA GAAGATAAAA   1500
146 AGTAGTATTC AAATGCACAT CAACGTCTCC ACTTGGAGGG CTTAAAGACG TTTCAACATA   1560
148 CAAACCGGGG AGTTTTGCCT GGAATGTTTC CTAAAATGTG TCCTGTAGCA CATAGGGTCC   1620
150 TCTTGTTCCT TAAAATCTAA TTACTTTTAG CCCAGTGCTC ATCCCACCTA TGGGGAGATG   1680
152 AGAGTGAAAA GGGAGCCTGA TTAATAATTA CACTAAGTCA ATAGGCATAG AGCCAGGACT   1740
154 GTTTGGGTAA ACTGGTCACT TTATCTTAAA CTAAATATAT CCAAAACTGA ACATGTACTT   1800
156 AGTTACTAAG TCTTTGACTT TATCTCATTC ATACCACTCA GCTTTATCCA GGCCACTTAT   1860
158 TTGACAGTAT TATTGCGAAA ACTTCCTAAC TGGTCTCCTT ATCATAGTCT TATCCCCTTT   1920
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160 TGAAACAAAA GAGACAGTTT CAAAATACAA ATATGATTTT TATTAGCTCC CTTTTGTTGT 1980
162 CTATAATAGT CCCAGAAGGA GTTATAAACT CCATTTAAAA AGTCTTTGAG ATGTGGCCCT 2040
164 TGCCAACTTT GCCAGGAATT CCCAATATCT AGTATTTTCT ACTATTAAAC TTTGTGCCTC 2100
166 TTCAAAACTG CATTTTCTCT CATTCCCTAA GTGTGCATTG TTTTCCCTTA CCGGTTGGTT 2160
168 TTTCCACCAC CTTTTACATT TTCCTGGAAC ACTATACCCT CCCTCTTCAT TTGGCCCACC 2220
170 TCTAATTTTC TTTCAGATCT CCATGAAGAT GTTACTTCCT CCAGGAAGCC TTATCTGACC 2280
172 CCTCCAAAGA TGTCATGAGT TCCTCTTTTC ATTCTACTAA TCACAGCATC CATCACACCA 2340
174 TGTTGTGATT ACTGATACTA TTGTCTGTTT CTCTGATTAG GCAGTAAGCT CAACAAGAGC 2400
176 TACATGGTGC CTGTCTCTTG TTGCTGATTA TTCCCATCCA AAAACAGTGC CTGGAATGCA 2460
178 GACTTAACAT TTTATTGAAT GAATAAATAA AACCCCATCT ATCGAGTGCT ACTTTGTGCA 2520
180 AGACCCGGTT CTGAGGCATT TATATTTATT GATTTATTTA ATTCTCATTT AACCATGAAG 2580
182 GAGGTACTAT CACTATCCTT ATTTTATAGT TGATAAAGAT AAAGCCCAGA GAAATGAATT 2640
184 AACTCACCCA AAGTCATGTA GCTAAGTCAC AGGGCAAAAA TTCAAACCAG TTCCCCAACT 2700
186 TTACGTGATT AATACTGTGC TATACTGCCT CTCTGATCAT ATGGCATGGA ATGCAGACAT 2760
188 CTGCTCCGTA AGGCAGAATA TGGAAGGAGA TTGGAGGATG ACACAAAACC AGCATAATAT 2820
190 CAGAGGAAAA GTCAAACAG GACCTGAACT GATAGAAAG TTGTTACTCC TGGTGTAGTC 2880
192 GCATCGACAT CTTGATGAAC TGGTGGCTGA CACAACATAC ATTGGCTTGA TGTGTACATA 2940
194 TTATTTGTAG TTGTGTGTGT ATTTTTATAT ATATATTTGT AATATTGAAA TAGTCATAAT 3000
196 TTACTAAAGG CCTACCATTT GCCAGGCATT TTTACATTTG TCCCCTCTAA TCTTTTGATG 3060
198 AGATGATCAG ATTGGATTAC TTGGCCTTGA AGATGATATA TCTACATCTA TATCTATATC 3120
200 TATATCTATA TCTATATCTA TATCTATATC TATATCTATA TATGTATATC AGAAAAGCTG 3180
202 AAATATGTTT TGTAAAGTTA TAAAGATTTC AGACTTTATA GAATCTGGGA TTTGCCAAAT 3240
204 GTAACCCCTT TCTCTACATT AAACCCATGT TGGAACAAAT ACATTTATTA TTCATTCATC 3300
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208 CATGCTTGGG CAAGCTTATA TAGTTTGCTT CATAAAACTC TATTTCAGTT CTTCATAACT 3420
210 AATACTTCAT GACTATTGCT TTTCAGGTAT TCCTTCATAA CAAATACTTT GGCTTTCATA 3480
212 TATTTGAGTA AAGTCCCCCT TGAGGAAGAG TAGAAGAACT GCACTTTGTA AATACTATCC 3540
214 TGGAATCCAA ACGGATAGAC AAGGATGGTG CTACCTCTTT CTGGAGAGTA CGTGAGCAAG 3600
216 GCCTGTTTTG TTAACATGTT CCTTAGGAGA CAAAACTTAG GAGAGACACG CATAGCAGAA 3660
218 AATGGACAAA AACTAACAAA TGAATGGGAA TTGTACTTGA TTAGCATTGA AGACCTTGTT 3720
220 TATACTATGA TAAATGTTTG TATTTGCTGG AAGTGCTACT GACGGTAAAC CCTTTTTGTT 3780
222 TAAATGTGTG CCCTAGTAGC TTGCAGTATG ATCTATTTTT TAAGTACTGT ACTTAGCTTA 3840
224 TTTAAAAATT TTATGTTTAA AATTGCATAG TGCTCTTTCA TTGAAGAAGT TTTGAGAGAG 3900
226 AGATAGAATT AAATTCACTT ATCTTACCAT CTAGAGAAAC CCAATGTTAA AACTTTGTTG 3960
228 TCCATTATTT CTGTCTTTTA TTCAACATTT TTTTTAGAGG GTGGGAGGAA TACAGAGGAG 4020
230 GTACAATGAT ACACAAATGA GAGCACTCTC CATGTATTGT TTTGTCCTGT TTTTCAGTTA 4080
232 ACAATATATT ATGAGCATAT TTCCATTTCA TTAAATATTC TTCCACAAAG TTATTTTGAT 4140
234 GGCTGTATAT CACCCTACTT TATGAATGTA CCATATTAAT TTATTTCCTG GTGTGGGTTA 4200
236 TTTGATTTTA TAATCTTACC TTTAGAATAA TGAAACACCT GTGAAGCTTT AGAAAATACT 4260
238 GGTGCCTGGG TCTCAACTCC ACAGATTCTG ATTTAACTGG TCTGGGTTAC AGACTAGGCA 4320
240 TTGGGAATTC AAAAAGTTCC CCCAGTGATT CTAATGTGTA GCCAAGATCG GGAACCCTTG 4380
242 TAGACAGGGA TGATAGGAGG TGAGCCACTC TTAGCATCCA TCATTTAGTA TTAACATCAT 4440
244 CATCTTGAGT TGCTAAGTGA ATGATGCACC TGACCCACTT TATAAAGACA CATGTGCAAA 4500
246 TAAAATTATT ATAGGACTTG GTTTATTAGG GCTTGTGCTC TAAGTTTTCT ATGTTAAGCC 4560
248 ATACATCGCA TACTAAATAC TTTAAAATGT ACCTTATTGA CATACATATT AAGTGAAAAG 4620
250 TGTTTCTGAG CTAAACAATG ACAGCATAAT TATCAAGCAA TGATAATTTG AAATGAATTT 4680
252 ATTATTCTGC AACTTAGGGA CAAGTCATCT CTCTGAATTT TTTGTACTTT GAGAGTATTT 4740
254 GTTATATTTG CAAGATGAAG AGTCTGAATT GGTCAGACAA TGTCTTGTGT GCCTGGCATA 4800
256 TGATAGGCAT TTAATAGTTT TAAAGAATTA ATGTATTTAG ATGAATTGCA TACCAAATCT 4860
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RAW SEQUENCE LISTING   DATE: 11/13/2000
PATENT APPLICATION: US/09/509,591   TIME: 10:31:59

Input Set : A:\Pto.amc
Output Set: N:\CRF3\11132000\I509591.raw

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258 GCTGTCTTTT CTTTATGGCT TCATTAACTT AATTTGAGAG AAATTAATTA TTCTGCAACT    4920
260 TAGGGACAAG TCATGTCTTT GAATATTCTG TAGTTTGAGG AGAATATTTG TTATATTTGC    4980
262 AAAATAAAAT AAGTTTGCAA GTTTTTTTTT TCTGCCCCAA AGAGCTCTGT GTCCTTGAAC    5040
264 ATAAAATACA AATAACCGCT ATGCTGTTAA TTATTGGCAA ATGTCCCATT TTCAACCTAA    5100
266 GGAAATACCA TAAAGTAACA GATATACCAA CAAAAGGTTA CTAGTTAACA GGCATTGCCT    5160
268 GAAAAGAGTA TAAAAGAATT TCAGCATGAT TTTCCATATT GTGCTTCCAC CACTGCCAAT    5220
270 AACA                                                                5224
272 (2) INFORMATION FOR SEQ ID NO: 3:
274      (i) SEQUENCE CHARACTERISTICS:
275           (A) LENGTH: 19 base pairs
276           (B) TYPE: nucleic acid
277           (C) STRANDEDNESS: single
278           (D) TOPOLOGY: linear
280      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
282 TCGTCTTCAA GAATTCTCA                                                 19
284 (2) INFORMATION FOR SEQ ID NO: 4:
286      (i) SEQUENCE CHARACTERISTICS:
287           (A) LENGTH: 20 base pairs
288           (B) TYPE: nucleic acid
289           (C) STRANDEDNESS: single
290           (D) TOPOLOGY: linear
292      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
294 TTTCAGTCAC CGGTGTCGGA                                                20
299 (2) INFORMATION FOR SEQ ID NO: 5:
301      (i) SEQUENCE CHARACTERISTICS:
302           (A) LENGTH: 20 base pairs
303           (B) TYPE: nucleic acid
304           (C) STRANDEDNESS: single
305           (D) TOPOLOGY: linear
307      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
309 GCATTCTCTA GACACAGGTG                                                20
311 (2) INFORMATION FOR SEQ ID NO: 6:
313      (i) SEQUENCE CHARACTERISTICS:
314           (A) LENGTH: 20 base pairs
315           (B) TYPE: nucleic acid
316           (C) STRANDEDNESS: single
317           (D) TOPOLOGY: linear
319      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
321 TCCGACACCG GTGACTGAAA                                                20
323 (2) INFORMATION FOR SEQ ID NO: 7:
325      (i) SEQUENCE CHARACTERISTICS:
326           (A) LENGTH: 21 base pairs
327           (B) TYPE: nucleic acid
328           (C) STRANDEDNESS: single
329           (D) TOPOLOGY: linear
331      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
333 GCCCACGGCC GCATTATATA C                                              21
335 (2) INFORMATION FOR SEQ ID NO: 8:
337      (i) SEQUENCE CHARACTERISTICS:
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338          (A) LENGTH: 21 base pairs
339          (B) TYPE: nucleic acid
340          (C) STRANDEDNESS: single
341          (D) TOPOLOGY: linear
343     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
345 GTATATAATG CGGCCGTGGG C                            21
348 (2) INFORMATION FOR SEQ ID NO: 9:
350     (i) SEQUENCE CHARACTERISTICS:
351          (A) LENGTH: 21 base pairs
352          (B) TYPE: nucleic acid
353          (C) STRANDEDNESS: single
354          (D) TOPOLOGY: linear
356     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
358 CCAGAAAATC CAGCAGGTAC C                            21
360 (2) INFORMATION FOR SEQ ID NO: 10:
362     (i) SEQUENCE CHARACTERISTICS:
363          (A) LENGTH: 29 base pairs
364          (B) TYPE: nucleic acid
365          (C) STRANDEDNESS: single
366          (D) TOPOLOGY: linear
368     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
370 GTGACCGGTG CATTGCTGTG AACTCTGTA                    29
372 (2) INFORMATION FOR SEQ ID NO: 11:
374     (i) SEQUENCE CHARACTERISTICS:
375          (A) LENGTH: 27 base pairs
376          (B) TYPE: nucleic acid
377          (C) STRANDEDNESS: single
378          (D) TOPOLOGY: linear
380     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
382 ATAAGTGGCC TGGATAAAGC TGAGTGG                      27
384 (2) INFORMATION FOR SEQ ID NO: 12:
386     (i) SEQUENCE CHARACTERISTICS:
387          (A) LENGTH: 28 base pairs
388          (B) TYPE: nucleic acid
389          (C) STRANDEDNESS: single
390          (D) TOPOLOGY: linear
392     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:
394 GTCACCGGTC TTTGTTATTG GCAGTGGT                     28
397 (2) INFORMATION FOR SEQ ID NO: 13:
399     (i) SEQUENCE CHARACTERISTICS:
400          (A) LENGTH: 30 base pairs
401          (B) TYPE: nucleic acid
402          (C) STRANDEDNESS: single
403          (D) TOPOLOGY: linear
405     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:
407 ATCCAGGCCA CTTATGAGCT CTGTGTCCTT                   30
409 (2) INFORMATION FOR SEQ ID NO: 14:
411     (i) SEQUENCE CHARACTERISTICS:
412          (A) LENGTH: 26 base pairs
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VERIFICATION SUMMARY   DATE: 11/13/2000
PATENT APPLICATION: US/09/509,591   TIME: 10:32:00

Input Set : **A:\Pto.amc**
Output Set: N:\CRF3\11132000\I509591.raw

L:30 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:31 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]

G.Lee

1632

```
RAW SEQUENCE LISTING             DATE: 11/06/2000
PATENT APPLICATION:  US/09/509,591   TIME: 12:25:12

Input Set : A:\#523643 v1 - 34802-20004.20 App. 09509591.txt
Output Set: N:\CRF3\11062000\I509591.raw
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Does Not Comply
Corrected Diskette Needed

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          SEQUENCE LISTING
 3 (1) GENERAL INFORMATION:
 5        (i) APPLICANT: Little, Andrew
 6                       Lamparski, Henry
 7                       Schuur, Eric
 8                       Henderson, Daniel
10       (ii) TITLE OF INVENTION: ADENOVIRUS VECTORS SPECIFIC FOR CELLS
11                        EXPRESSING ALPHA-FETOPROTEIN AND METHODS OF USE THEREOF
13      (iii) NUMBER OF SEQUENCES: 23
15       (iv) CORRESPONDENCE ADDRESS:
16            (A) ADDRESSEE: MORRISON & FOERSTER
17            (B) STREET: 755 PAGE MILL ROAD
18            (C) CITY: PALO ALTO
19            (D) STATE: CA
20            (E) COUNTRY: USA
21            (F) ZIP: 94304-1018
23        (v) COMPUTER READABLE FORM:
24            (A) MEDIUM TYPE: Floppy disk
25            (B) COMPUTER: IBM PC compatible
26            (C) OPERATING SYSTEM: PC-DOS/MS-DOS
27            (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
29       (vi) CURRENT APPLICATION DATA:
C--> 30       (A) APPLICATION NUMBER: US/09/509,591
C--> 31       (B) FILING DATE: 02-Jun-2000
32            (C) CLASSIFICATION:
34      (vii) PRIOR APPLICATION DATA:
35            (A) APPLICATION NUMBER: PCT/US98/04084
36            (B) FILING DATE: 03-MAR-1998
38     (viii) ATTORNEY/AGENT INFORMATION:
39            (A) NAME: POLIZZI, CATHERINE M.
40            (B) REGISTRATION NUMBER: 40,130
41            (C) REFERENCE/DOCKET NUMBER: 348022000420
43       (ix) TELECOMMUNICATION INFORMATION:
44            (A) TELEPHONE: (650) 813-5600
45            (B) TELEFAX: (650) 494-0792
46            (C) TELEX: 706141 MRSNFOERS SFO


ERRORED SEQUENCES

549 (2) INFORMATION FOR SEQ ID NO: 23:
551       (i) SEQUENCE CHARACTERISTICS:
552           (A) LENGTH: 101 amino acids
553           (B) TYPE: amino acid
554           (D) TOPOLOGY: linear
556      (ii) MOLECULE TYPE: protein
558      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 23:
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RAW SEQUENCE LISTING   DATE: 11/06/2000
PATENT APPLICATION: US/09/509,591   TIME: 12:25:12

Input Set : A:\#523643 v1 - 34802-20004.20 App. 09509591.txt
Output Set: N:\CRF3\11062000\I509591.raw

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561   1               5                  10             15
563 Ala Thr Gly Leu Thr Ser Ala Leu Asn Leu Pro Gln Val His Ala Phe
564              20                  25             30
566 Val Asn Asp Trp Ala Ser Leu Asp Met Trp Trp Phe Ser Ile Ala Leu
567          35                  40              45
569 Met Phe Val Cys Leu Ile Ile Met Trp Leu Ile Cys Cys Leu Lys Arg
570          50                  55              60
572 Arg Arg Ala Arg Pro Pro Ile Tyr Arg Pro Ile Ile Val Leu Asn Pro
573  65                  70                  75             80
575 His Asn Glu Lys Ile His Arg Leu Asp Gly Leu Lys Pro Cys Ser Leu
576                  85                  90              95
578 Leu Leu Gln Tyr Asp
579                 100
E--> 585  1
E--> 588  1
E--> 589  |||
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VERIFICATION SUMMARY　　　　　　　　DATE: 11/06/2000
PATENT APPLICATION:　US/09/509,591　　　　TIME: 12:25:13

Input Set : A:\#523643 v1 - 34802-20004.20 App. 09509591.txt
Output Set: N:\CRF3\11062000\I509591.raw

L:30 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:31 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
L:585 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:23
M:332 Repeated in SeqNo=23
L:589 M:330 E: (2) Invalid Amino Acid Designator, 1
L:589 M:203 E: No. of Seq. differs, LENGTH:Input:101 Found:102 SEQ:23